

INFORMATION DISCLOSURE CITATION

Attorney Docket No.: GC626-3	Serial No.: 09/932,279
Applicant: Estell et al.	RECEIVED
Filing Date: August 17, 2001	
Page 2 of 2	Group: 1645 1631
	Date of this Submission: January 3, 2002

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Examiner	Date Considered
<i>Arden Mansley</i>	<i>1-9-04</i>

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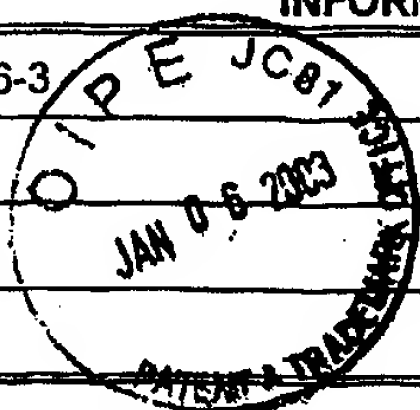
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Examiner's	
Initials	Author, Title, Date, Pertinent Pages, etc.
	*Allen, C., "Sequencing of Proteins and Peptides," 2nd edn. Elsevier, Amsterdam -- Book
AM	*Altschul et al., "Basic Local Alignment Search Tool," J. Mol. Biol., V. 215, 1990, pp. 403-410.
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	*Corthals et al., "Identification of Proteins by Mass Spectrometry," Ed. Rabilloud, T. Springer, New York, pp. 197-231, 1999.
	*Christianson et al., "Peptide Mapping of Subtilisins as a Practical tool for Locating Protein Sequence Errors during Extensive Protein Engineering Projects," Anal. Biochem. 1994, Vol. 223, pp. 119-129.
	*Deutscher MP, "Guide to Protein purification," Academic Press, New York. -- Book
AM	*George et al., "PIR-International Protein Sequence Database", Methods in Enzymology, V. 266, pp. 41-59 (1996).
	*Goddette et al., "The Crystal Structure of the Bacillus lentus Alkaline Protease, Subtilisin BL, at 1.4 A Resolution", J. Mol. Biol. V. 228, pp. 580-595 (1992).
	*Goeddel et al., "Synthesis of Human Fibroblast Interferon by E. coli," Nucl. Acids Res., Vol. 8, pp. 4057-4074, 1980.
	*Guermant et al., "Under Proper Control, Oxidation of Proteins with Known Chemical Structure Provides an Accurate and Absolute Method for the Determination of Their Molar Concentration", Anal. Biochem. V. 277, pp. 46-57 (2000).
	*Gygi et al., "Quantitative analysis of complex protein mixtures using isotope-coded affinity tags," Nature Biotech. V. 17, pp. 994-999 (1999).
	*Hale & Marham, The Harper Collins Dictionary of Biology, Harper Perennial, NY (1991) -- Book
	*Hancock ES, "New Methods in Peptide Mapping for the Characterization of Proteins", CRC Press, Boca Raton -- Book
AM	*Hsia et al., "Active-Site Titration of Serine Proteases Using a Fluoride Ion Selective Electrode and Sulfonyl Fluoride Inhibitors," Analytical Biochem., V. 242, pp. 221-227 (1996).
	*Janson et al., "Protein Purification," 2nd edn., Wiley-Liss, New York -- Book
	*Kellner et al., "Microcharacterization of Proteins", 2nd edn., Wiley-VCH, Weinheim -- Book
AM	*Khan et al., "EMBL Data Library", Methods in Enzymology, V. 183, pp. 23-31 (1990).
	Karlin et al., "Application and Statistics for Multiple High-Scoring Segments in Molecular Sequence," Proc. Natl. Acad. Sci. USA, V. 90, pp. 5873-5877 (1993).
	*Karlin et al., "Methods for assessing the statistical significance of molecular sequence features by using general scoring schemes," Proc. Natl. Acad. Sci. USA, V. 87, pp. 2264-2268 (1990).

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John Marable	1-9-04

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